RAW SEQUENCE LISTING DATE: 05/23/2001
PATENT APPLICATION: US/09/831,233 TIME: 16:09:29

Input Set : A:\BB1129 seq lst.txt
Output Set: C:\CRF3\05232001\I831233.raw

3 <110> APPLICANT: E. I. du Pont de Nemours and Company
5 <120> TITLE OF INVENTION: Plant Glutamine Amidotransferase Homologs
7 <130> FILE REFERENCE: BB1129
C--> 9 <140> CURRENT APPLICATION NUMBER: US/09/831,233
C--> 10 <141> CURRENT FILING DATE: 2001-05-04
12 <150> PRIOR APPLICATION NUMBER: 60/107,275
13 <151> PRIOR FILING DATE: 1998-11-05
15 <160> NUMBER OF SEQ ID NOS: 15
17 <170> SOFTWARE: Microsoft Office 97
19 <210> SEQ ID NO: 1
20 <211> LENGTH: 1956
21 <212> TYPE: DNA
22 <213> ORGANISM: Impatiens balsamia
24 <400> SEQUENCE: 1

ENTERED
See p.5

25 gcacgagaag gaactggcaa gtaccaagcc acagaatggg ttcagaattc gcgccgcctt 60
26 ggctggtgca ggcggagatt ctgtggtgac tttacttgat tacggtgctg gaaatgttag 120
27 gagtgtgagg aacgccatcc gcacacttgg atttgatatc aaagatgtgc aaaagccaga 180
28 ggatattcta aatgctaagc gccttatctt tcctggcgtt ggggcctttg cacctgcaat 240
29 ggatgttctt attcgtaaag ggctggctga agcactctgt acttacattc agaatgatcg 300
30 acctttcctg ggtatatgcc tgggattgca gctactcttt gagtcaagtg aagaaaatgg 360
31 tccaattcaa ggtcttggct tgattcctgg acgggttggg cgttttgaat catccaatgg 420
32 tttaagggtg ccacatattg gatggcatgc cttggatata aaggaagggt cagcaatttt 480
33 agatgatgtg gggaatcaac atgtgtattt tgttcactca tatcgagcca atgccgagga 540
34 caacaaagag tggatttcat ctacatgcag ctatggtgac gattttattg catccattca 600
35 gaagggaaat gttcatgcag tccaatttca tcccgagaag agtggaggtg ttggactttc 660
36 catattgaga agatttttga atgctgattc ctttaacaac aaaagacaga agccaatgaa 720
37 tggaaaggct tctaaacttg caaagagagt aattgcttgc cttgatgtga gggcaaatga 780
38 taatggggat cttgttgtaa ccaagggaga ccaatatgat gtgagagaac gtacagaaga 840
39 gaatgaggtc agaaaccttg gcaagcctgt tgaacttgct gggcagtatt atttagacgg 900
40 tgctgatgag gtcagcttct taaacattac tggtttccgg gacttccctc taggcgatct 960
41 acccatgcta caggtcttgc aacgcgcatc tgaaaacgtt tttgtgccat taactgtcgg 1020
42 gggtggcatc agggatttta ctgatgcaaa tggaaggtat tattctagtc tagaagtggc 1080
43 ttcagagtat ttcagatcgg gcgccgataa ggtttcgatc ggaagtgatg cagtttacac 1140
44 tgctgaggaa tatattaaaa ccggagtgaa gacaggaaag agcagcatag agcagatatc 1200
45 tacagtatat ggtaaccagg cagtggttgt aagcattgat cctcgccgag tttacttgag 1260
46 aaaacccgat gaagtagaat ttaaagccat caaagtaagc catccaggtc caaacggtga 1320
47 ggaatatgcc tggtatcagt gcactgttaa tggtggacga gaagggagac ccatcggagc 1380
48 ttatgaacta gctaaggctg ttgaggaact tggagctgga gaaatattat tgaactgcat 1440
49 tgattgtgat ggtcaaggaa aaggattcga tatagatctg atcaagctaa tatccgatgc 1500
50 tgtgaacatt cctgttatcg caagcagcgg tgcaggagtc gctgatcact tctccgaagt 1560
51 ctttaatgaa accaacgcat ctgctgccct tgcagctggc attttccatc gcaaagaggt 1620
52 tccaattaag gctgttaaag agcacttgtt gaaggaaggg attgaagtta gattgtaagg 1680
53 cgagaatcac ttggaagaaa tttcatcttg aagttcaatt ttgttacaca agagatttcc 1740
54 ttctttcttg gcctatgtga tatttattta tttatgtttt gctattgaat tattgttatt 1800
55 attattttgg catttgttat ttgaatagat ttgagttttt agaccttggt gtgtcctgtt 1860
56 tatctctagg ccatgttttg tggattatat acaagtgtga aattaaataa ataaatcgta 1920
57 tgaatttatg cttttaaaaa aaaaaaaaaa aaaaaa 1956

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PATENT APPLICATION: **US/09/831,233** TIME: 16:09:29

Input Set : **A:\BB1129 seq lst.txt**
Output Set: **C:\CRF3\05232001\I831233.raw**

59 <210> SEQ ID NO: 2
60 <211> LENGTH: 558
61 <212> TYPE: PRT
62 <213> ORGANISM: Impatiens balsamia
64 <400> SEQUENCE: 2
65 His Glu Lys Glu Leu Ala Ser Thr Lys Pro Gln Asn Gly Phe Arg Ile
66 1 5 10 15
68 Arg Ala Ala Leu Ala Gly Ala Gly Gly Asp Ser Val Val Thr Leu Leu
69 20 25 30
71 Asp Tyr Gly Ala Gly Asn Val Arg Ser Val Arg Asn Ala Ile Arg Thr
72 35 40 45
74 Leu Gly Phe Asp Ile Lys Asp Val Gln Lys Pro Glu Asp Ile Leu Asn
75 50 55 60
77 Ala Lys Arg Leu Ile Phe Pro Gly Val Gly Ala Phe Ala Pro Ala Met
78 65 70 75 80
80 Asp Val Leu Ile Arg Lys Gly Leu Ala Glu Ala Leu Cys Thr Tyr Ile
81 85 90 95
83 Gln Asn Asp Arg Pro Phe Leu Gly Ile Cys Leu Gly Leu Gln Leu Leu
84 100 105 110
86 Phe Glu Ser Ser Glu Glu Asn Gly Pro Ile Gln Gly Leu Gly Leu Ile
87 115 120 125
89 Pro Gly Arg Val Gly Arg Phe Glu Ser Ser Asn Gly Leu Arg Val Pro
90 130 135 140
92 His Ile Gly Trp His Ala Leu Asp Ile Lys Glu Gly Ser Ala Ile Leu
93 145 150 155 160
95 Asp Asp Val Gly Asn Gln His Val Tyr Phe Val His Ser Tyr Arg Ala
96 165 170 175
98 Asn Ala Glu Asp Asn Lys Glu Trp Ile Ser Ser Thr Cys Ser Tyr Gly
99 180 185 190
101 Asp Asp Phe Ile Ala Ser Ile Gln Lys Gly Asn Val His Ala Val Gln
102 195 200 205
104 Phe His Pro Glu Lys Ser Gly Gly Val Gly Leu Ser Ile Leu Arg Arg
105 210 215 220
107 Phe Leu Asn Ala Asp Ser Phe Asn Asn Lys Arg Gln Lys Pro Met Asn
108 225 230 235 240
110 Gly Lys Ala Ser Lys Leu Ala Lys Arg Val Ile Ala Cys Leu Asp Val
111 245 250 255
113 Arg Ala Asn Asp Asn Gly Asp Leu Val Val Thr Lys Gly Asp Gln Tyr
114 260 265 270
116 Asp Val Arg Glu Arg Thr Glu Glu Asn Glu Val Arg Asn Leu Gly Lys
117 275 280 285
119 Pro Val Glu Leu Ala Gly Gln Tyr Tyr Leu Asp Gly Ala Asp Glu Val
120 290 295 300
122 Ser Phe Leu Asn Ile Thr Gly Phe Arg Asp Phe Pro Leu Gly Asp Leu
123 305 310 315 320
125 Pro Met Leu Gln Val Leu Gln Arg Ala Ser Glu Asn Val Phe Val Pro
126 325 330 335
128 Leu Thr Val Gly Gly Gly Ile Arg Asp Phe Thr Asp Ala Asn Gly Arg
129 340 345 350

RAW SEQUENCE LISTING DATE: 05/23/2001
PATENT APPLICATION: US/09/831,233 TIME: 16:09:29

Input Set : A:\BB1129 seq lst.txt
Output Set: C:\CRF3\05232001\I831233.raw

```
131 Tyr Tyr Ser Ser Leu Glu Val Ala Ser Glu Tyr Phe Arg Ser Gly Ala
132         355                 360                 365
134 Asp Lys Val Ser Ile Gly Ser Asp Ala Val Tyr Thr Ala Glu Glu Tyr
135     370                 375                 380
137 Ile Lys Thr Gly Val Lys Thr Gly Lys Ser Ser Ile Glu Gln Ile Ser
138 385                 390                 395                 400
140 Thr Val Tyr Gly Asn Gln Ala Val Val Val Ser Ile Asp Pro Arg Arg
141                 405                 410                 415
143 Val Tyr Leu Arg Lys Pro Asp Glu Val Glu Phe Lys Ala Ile Lys Val
144             420                 425                 430
146 Ser His Pro Gly Pro Asn Gly Glu Glu Tyr Ala Trp Tyr Gln Cys Thr
147         435                 440                 445
149 Val Asn Gly Gly Arg Glu Gly Arg Pro Ile Gly Ala Tyr Glu Leu Ala
150     450                 455                 460
152 Lys Ala Val Glu Glu Leu Gly Ala Gly Glu Ile Leu Leu Asn Cys Ile
153 465                 470                 475                 480
155 Asp Cys Asp Gly Gln Gly Lys Gly Phe Asp Ile Asp Leu Ile Lys Leu
156                 485                 490                 495
158 Ile Ser Asp Ala Val Asn Ile Pro Val Ile Ala Ser Ser Gly Ala Gly
159             500                 505                 510
161 Val Ala Asp His Phe Ser Glu Val Phe Asn Glu Thr Asn Ala Ser Ala
162         515                 520                 525
164 Ala Leu Ala Ala Gly Ile Phe His Arg Lys Glu Val Pro Ile Lys Ala
165     530                 535                 540
167 Val Lys Glu His Leu Leu Lys Glu Gly Ile Glu Val Arg Leu
168 545                 550                 555
170 <210> SEQ ID NO: 3
171 <211> LENGTH: 587
172 <212> TYPE: DNA
173 <213> ORGANISM: Zea mays
175 <220> FEATURE:
176 <221> NAME/KEY: unsure
177 <222> LOCATION: (495)
179 <220> FEATURE:
180 <221> NAME/KEY: unsure
181 <222> LOCATION: (518)
183 <220> FEATURE:
184 <221> NAME/KEY: unsure
185 <222> LOCATION: (577)
187 <220> FEATURE:
188 <221> NAME/KEY: unsure
189 <222> LOCATION: (582)
191 <400> SEQUENCE: 3
192 aagaaaaggc ccgctagggc gccgagacag cggaacgttc tctgagtttg agcacgatct  60
193 ccccgggccc cggcgccgcc gtacgtcccc cttcggcgtc gccagccgcc tcctggctcc 120
194 ggctccttca tcgcctgctc cagcgtgcct gcgtgacata agcgtcgatt gattggcgag 180
195 aaaggggacg aatgcagccg ccgttgcagg cgcagggagc aatggctaac gtcgccgcta 240
196 tcctcaccgt cccctgctcc gcgggccgcc gcccgaagcg gagcaaccag ccccgcggat 300
197 gcggctccgt ctccgtctcc gtctccgtcc gtgcgtcctc cggcgcaaac acggtgactc 360
```

RAW SEQUENCE LISTING DATE: 05/23/2001
PATENT APPLICATION: **US/09/831,233** TIME: 16:09:29

Input Set : **A:\BB1129 seq lst.txt**
Output Set: **C:\CRF3\05232001\I831233.raw**

```
     198 tgctggacta cggcgcgggg aacgtacgca gcgtgcgcaa cgcaattcgc tacctcggct 420
     199 tcgacatccg cgacgtgcag agcccggarg acatcgtcgc cggcggaayg ggtcgtcttt 480
W--> 200 cccggtgtcg gcgcnttcgg ctccgccatg gacgtccnca ccaggacggg catgccaacg 540
W--> 201 cactccgtga gtacatccaa agggaacgcc ccttccnagg cnctgcc               587
     203 <210> SEQ ID NO: 4
     204 <211> LENGTH: 130
     205 <212> TYPE: PRT
     206 <213> ORGANISM: Zea mays
     208 <220> FEATURE:
     209 <221> NAME/KEY: UNSURE
     210 <222> LOCATION: (90)
     212 <220> FEATURE:
     213 <221> NAME/KEY: UNSURE
     214 <222> LOCATION: (93)
     216 <220> FEATURE:
     217 <221> NAME/KEY: UNSURE
     218 <222> LOCATION: (109)
     220 <220> FEATURE:
     221 <221> NAME/KEY: UNSURE
     222 <222> LOCATION: (115)
     224 <220> FEATURE:
     225 <221> NAME/KEY: UNSURE
     226 <222> LOCATION: (129)
     228 <400> SEQUENCE: 4
     229 Met Gln Pro Pro Leu Gln Ala Gln Gly Ala Met Ala Asn Val Ala Ala
     230   1               5                  10                  15
     232 Ile Leu Thr Val Pro Cys Ser Ala Gly Arg Arg Pro Lys Arg Ser Asn
     233              20                  25                  30
     235 Gln Pro Arg Gly Cys Gly Ser Val Ser Val Ser Val Ser Val Arg Ala
     236          35                  40                  45
     238 Ser Ser Gly Ala Asn Thr Val Thr Leu Leu Asp Tyr Gly Ala Gly Asn
     239      50                  55                  60
     241 Val Arg Ser Val Arg Asn Ala Ile Arg Tyr Leu Gly Phe Asp Ile Arg
     242  65                  70                  75                  80
W--> 244 Asp Val Gln Ser Pro Glu Asp Ile Val Xaa Ala Glu Xaa Val Val Phe
     245                  85                  90                  95
W--> 247 Pro Gly Val Gly Ala Phe Gly Ser Ala Met Asp Val Xaa Thr Arg Thr
     248              100                 105                 110
W--> 250 Gly Met Xaa Asn Ala Leu Arg Glu Tyr Ile Gln Arg Glu Arg Pro Phe
     251          115                 120                 125
W--> 253 Xaa Gly
     254      130
     256 <210> SEQ ID NO: 5
     257 <211> LENGTH: 1084
     258 <212> TYPE: DNA
     259 <213> ORGANISM: Zea mays
     261 <400> SEQUENCE: 5
     262 actagtggta acaaaaggcg atcaatatga tgtaagagat catactagca gcaaagaggt   60
     263 aagaaacctt ggcaagccag tcgatttagc aagccagtac tacatagacg gtgctgatga  120
```

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PATENT APPLICATION: **US/09/831,233** TIME: 16:09:29

Input Set : **A:\BB1129 seq lst.txt**
Output Set: **C:\CRF3\05232001\I831233.raw**

```
264 ggtcagcttc ttgaatataa ctggtttccg tgactttcca ttgggtgatt tgccaatgct  180
265 agaggtactg cgttgtgcct ctgaaaaggt ttttgtgcca cttacagttg gtgggggcat  240
266 acgagacttc acagatgcaa atggaagata ctactcaagt ttggaggtag catcagaata  300
267 tttcaggtcc ggtgctgaca aaatttcaat tggaagtgat gctgtttatg ctgctgaagc  360
268 ctttttacag actggtgtaa agacagggaa aagcagcttg gagcaaatct ctagagtata  420
269 tggcaatcag gctgtagttg tcagtattga tcctcgacgg gtatatgtca aaagtcaaga  480
270 agatgtgcca tttaaaactg taaaggtgtc cactaaaggt ccatcgggag aagaatatgc  540
271 atggtaccag tgcacagtga atggtggacg tgatagccga gctataggag catatgaact  600
272 agcgaaagct gtggaagaat tgggcgcagg agaaatactt cttaactgca ttgattgtga  660
273 tggccaaggt tgtggatttg acatagattt ggttaaaatg gtttctgatg ctgtgacaat  720
274 ccctgtcatt gcgagcagtg gtgctggagc tgttcaacat ttttctgaaa tttttgagaa  780
275 aacaaatgct tctgctgccc ttgctgctgg cattttccac cggaaagagg ttcctatact  840
276 agcagtgaaa gagcatctgg tcaatgctgg tgtggaggtc agggtgtaac agggagatcc  900
277 ttcggtttat tgaaatattc ttgtttgatg tcacaactgc tatcagttct gtttctctga  960
278 tgtcgcaact gctatcagat ctgttggtgg cagctggcag tgcataggcc cctgtcgaga 1020
279 actgcagttt ggtaataaat taataatgtg atgcttaaca gattaaaaaa aaaaaaaaaa 1080
280 aaaa                                                             1084
282 <210> SEQ ID NO: 6
283 <211> LENGTH: 295
284 <212> TYPE: PRT
285 <213> ORGANISM: Zea mays
287 <400> SEQUENCE: 6
288 Leu Val Val Thr Lys Gly Asp Gln Tyr Asp Val Arg Asp His Thr Ser
289   1               5                  10                  15
291 Ser Lys Glu Val Arg Asn Leu Gly Lys Pro Val Asp Leu Ala Ser Gln
292              20                  25                  30
294 Tyr Tyr Ile Asp Gly Ala Asp Glu Val Ser Phe Leu Asn Ile Thr Gly
295          35                  40                  45
297 Phe Arg Asp Phe Pro Leu Gly Asp Leu Pro Met Leu Glu Val Leu Arg
298      50                  55                  60
300 Cys Ala Ser Glu Lys Val Phe Val Pro Leu Thr Val Gly Gly Gly Ile
301  65                  70                  75                  80
303 Arg Asp Phe Thr Asp Ala Asn Gly Arg Tyr Tyr Ser Ser Leu Glu Val
304                  85                  90                  95
306 Ala Ser Glu Tyr Phe Arg Ser Gly Ala Asp Lys Ile Ser Ile Gly Ser
307             100                 105                 110
309 Asp Ala Val Tyr Ala Ala Glu Ala Phe Leu Gln Thr Gly Val Lys Thr
310         115                 120                 125
312 Gly Lys Ser Ser Leu Glu Gln Ile Ser Arg Val Tyr Gly Asn Gln Ala
313     130                 135                 140
315 Val Val Val Ser Ile Asp Pro Arg Arg Val Tyr Val Lys Ser Gln Glu
316 145                 150                 155                 160
318 Asp Val Pro Phe Lys Thr Val Lys Val Ser Thr Lys Gly Pro Ser Gly
319                 165                 170                 175
321 Glu Glu Tyr Ala Trp Tyr Gln Cys Thr Val Asn Gly Gly Arg Asp Ser
322             180                 185                 190
324 Arg Ala Ile Gly Ala Tyr Glu Leu Ala Lys Ala Val Glu Glu Leu Gly
325         195                 200                 205
327 Ala Gly Glu Ile Leu Leu Asn Cys Ile Asp Cys Asp Gly Gln Gly Cys
```

F.Y.I. →

Please Note:
Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

VERIFICATION SUMMARY DATE: 05/23/2001
PATENT APPLICATION: **US/09/831,233** TIME: 16:09:30

Input Set : **A:\BB1129 seq lst.txt**
Output Set: **C:\CRF3\05232001\I831233.raw**

L:9 M:270 C: Current Application Number differs, Replaced Application Number
L:10 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:200 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:3
L:200 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3
L:201 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:3
L:201 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3
L:244 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:4
L:244 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4
L:247 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:4
L:247 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4
L:250 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:4
L:250 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4
L:253 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:4
L:253 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4
L:451 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:7
L:451 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7
L:453 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:7
L:453 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7
L:454 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:7
L:454 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7
L:455 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:7
L:455 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7
L:456 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:7
L:456 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7
L:457 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:7
L:457 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7
L:458 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:7
L:458 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7
L:502 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:8
L:502 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8
L:505 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:8
L:505 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8
L:508 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:8
L:508 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8
L:511 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:8
L:511 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8
L:585 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:9
L:585 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9
L:587 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:9
L:587 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9
L:588 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:9
L:588 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9
L:589 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:9
L:589 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9
L:590 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:9
L:590 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9
L:591 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:9
L:591 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9

VERIFICATION SUMMARY DATE: 05/23/2001
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Input Set : **A:\BB1129 seq lst.txt**
Output Set: **C:\CRF3\05232001\I831233.raw**

L:615 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:10
L:615 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10
L:663 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:13
L:663 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13
L:697 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:14
L:697 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:14